Human GH Signal Peptide Genomic Sequence

G GTAAGCGC CCCAAGCTIGCCACC AIG GCT ACA

ATG GCT ACA G GTAAGCGCCC CTAAAATCCC TTTGGGCACA Met Ala Thr G

CTGGAGGGAT GGAGAGAA AAACAAACAG CTCCTGGAGC AGGGAGAGTG ATTTGGCCAA TCTCAGAAAG ATGTGTCCTG AGGGGAGAGG CAGCGACCTG TAGATGGGAC GGGGGCACTA ACCCTCAGGT GCTCTCCGGC ICCCICIGIT GCCCICIGGI ITCICCCCAG TIGGGGCIIC IGAAIGIGAG IAICGCCAIG IAAGCCCAGI CICCIGGICC CIGGCCICII

ly ser Arg Thr Ser Leu Leu Leu Ala Phe Gly Leu Leu Cys GC TCC CGG ACG TCC CTG CTC CTG GCT TTT GGC CTG CTC TGC

GAC GCT TGG CIT CAA GAG GGC AGT GCC AIG GCT TIA

signal peptide cleavage site 1

Pro Trp Leu Gln Glu Gly Ser Ala CCC TGG CTT CAA GAG GGC AGT GCC

Leu

Fig. 1A

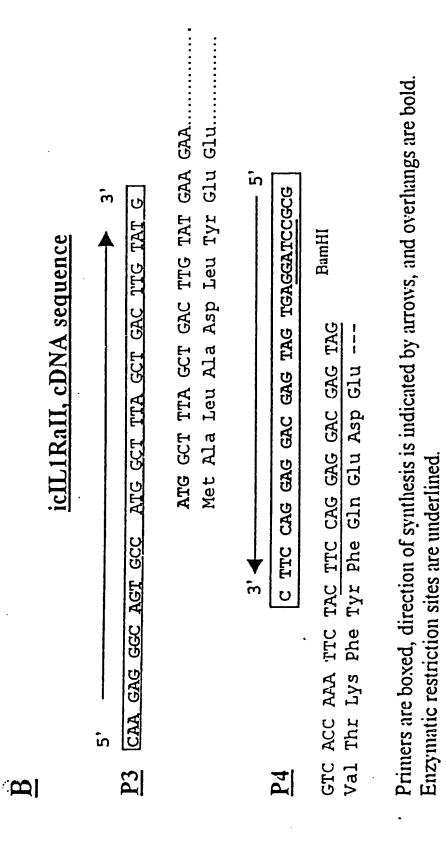
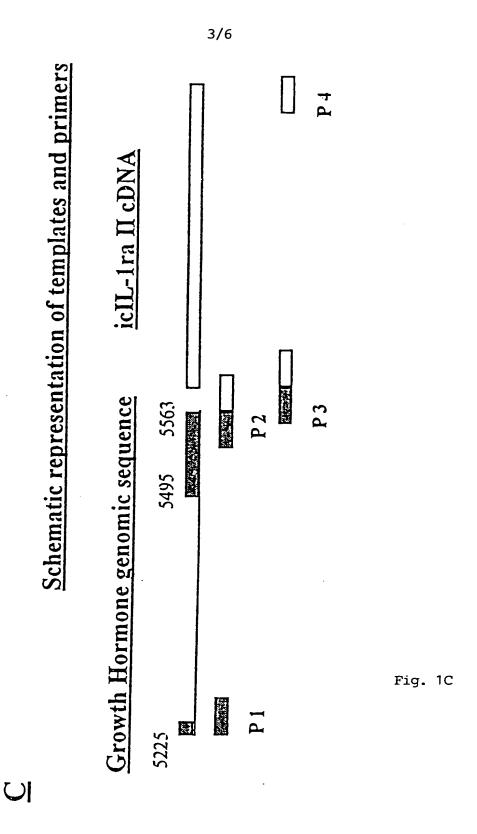


Fig. 1B



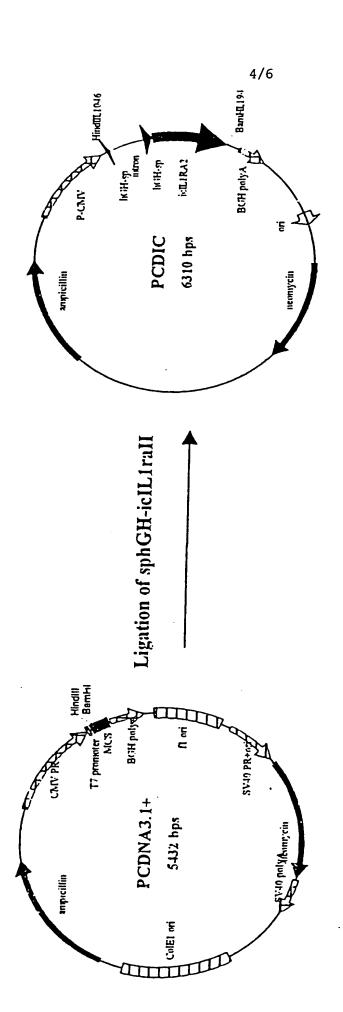
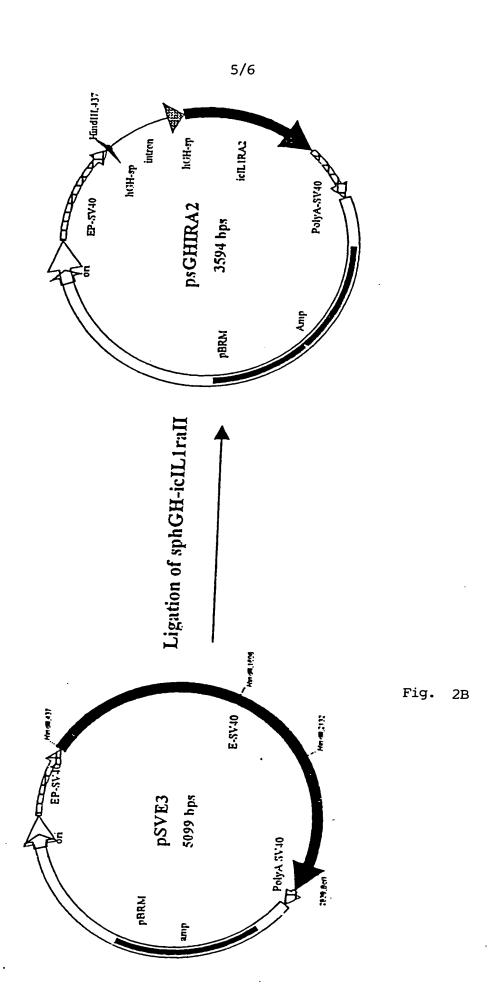


Fig. 2A



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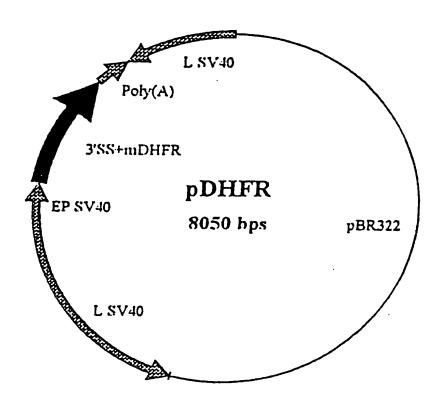


Fig. 2C

Human GH Signal Peptide Genomic Sequence

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ATG GCT ACA G GTAAGCGCCC CTAAAATCCC TTTGGGCACA G GIAAGCGC PI CCCAAGCTIGCCACC AIG GCT ACA

HindIII

Met Ala Thr G

TGAATGTGAG TATCGCCATG TAAGCCCAGT ATTTGGCCAA TCTCAGAAAG AGGGGAGAGG CAGCGACCTG TAGATGGGAC GGGGGCACTA ACCCTCAGGT CIGGAGGGAI GGAGAGAA AAACAAACAG CICCIGGAGC CIGGCCICIT GCICICGGC ICCCICIGIT GCCCICIGGI IICICCCCAG ATGTGTCCTG TIGGGGCTIC

ly ser Arg Thr ser Leu Leu Leu Ala Phe Gly Leu Leu Cys GC TCC CGG ACG TCC CTG CTG GCT TTT GGC CTG CTC TGC

GCC ATG GCA GAG GGC AGT AAA C TEG CIT. G TCA GCT

Leu Gln Glu Gly Ser Ala CCC TGG CIT CAA GAG GGC AGT GCC Pro Trp

signal peptide cleavage site

1A Fig.

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icIL1RaII, cDNA sequence

GAG GGC AGT GCC

ATG GCT TTA GCT GAC TTG TAT GAA GAA.... Met Ala Leu Ala Asp Leu Tyr Glu Glu....

3, 4 GAT CET CAC TAC TCG TCCTGGAAGS (S) CO TITE CAG GAG GAG GAG TAG TGAGGATCCGGG

BamHI

GTC ACC AAA TTC TAC TTC CAG GAG GAC GAG TAG Val Thr Lys Phe Tyr Phe Gln Glu Asp Glu --- Primers are boxed, direction of synthesis is indicated by arrows, and overhangs are bold. Enzymatic restriction sites are underlined.

Fig. 1B